# How to Utilize Side Information: Decomposed Gaussian Process Regression

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#### Problem Background

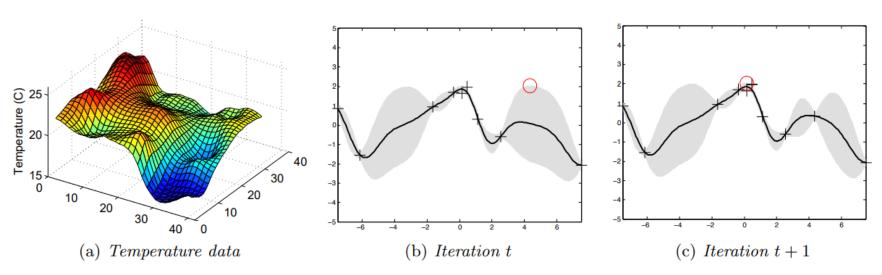
- ► Goal: we want to minimize the total Tuberculosis infected population in the following 25 years.
- ▶ Input: policy  $\nu \in \mathbb{R}^n$ , n = 110
  - Where  $v_i$  refers to what proportion of people in age i would catch the information about Tuberculosis (so that they will seek a treatment and get cured.)
- ▶ Budget constraint:  $\sum v_i \leq B$

# Problem Background

- ▶ What we have:
  - A very expensive simulation model (or a field test) that can simulate the interaction between people. Let's call this function *f*
  - $\blacktriangleright f(v)$  represents the outcome by applying policy v.
  - ▶ But it takes 30 minutes to run one f(v) ...

# Gaussian Process Upper-confidence Based

$$\mu_T(x) = k_T(x)^T (K_T + \sigma^2 I)^{-1} y_T \qquad k_T(x) = [k(x_1, x), \dots, k(x_m, x)]^T \\ k_T(x, x') = k(x, x') - k_T(x)^T (K_T + \sigma^2 I)^{-1} k_T(x') \qquad K_T = [k(x, x')]_{x, x' \in A_T} \in S_{++}^n \\ \sigma_T^2(x) = k_T(x, x)$$



Gaussian Process Optimization in the Bandit Setting: No Regret and Experimental Design, 2010

# In our problem

- We can use Gaussian process regression to approximate our  $f(v), v \in \mathbb{R}^n$
- ▶ But we don't know the kernel k(x, x')...

#### Markov property

- When we run a simulation, it will give us the total infected population of age i at time t, which is denoted by  $I_i^t$ .
- Also the healthy population  $S_i^t$  and latent TB population  $E_i^t$
- Let  $x_t = [I_i^t, S_i^t, E_i^t]$  be the collection of all variables at time t.

# Markov property

- memoryless property
  - $\triangleright E[x_{t+1}|x_t, x_{t-1}, ..., x_1, v] = E[x_{t+1}|x_t, v]$
- So we can write down the variables at time t+1 as  $h_t(x_t, v)$

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- ► Motivated by the SEIS population model:
  - $I_{i+1}^{t+1} = I_i^t (1 d_i^t) (1 \nu_i) + E_i^t \alpha_i^t$
  - $E_{i+1}^{t+1} = E_i^t (1 \mu_i^t) (1 \alpha_i^t) + S_i^t (1 \mu_i^t) \sum_{k=1}^n \beta_{ik} \frac{I_k^t}{I_k^t + E_k^t + S_k^t}$
  - $\approx E_i^t (1 \mu_i^t) (1 \alpha_i^t) + constant (1 \mu_i^t) \sum_{k=1}^n \beta_{ik} \frac{I_k^t}{constant}$

#### Linear approximation

- $> x_{t+1} = h_t(x_t, v) = A_t(v)x_t + f_t(v)$ 
  - Where  $A_t(v) \in \mathbb{R}^{n \times n}$  is a matrix of linear function in terms of v
  - ▶ And  $f_t(v)$  is the difference (error) of  $x_{t+1}$  and  $A_t(v)x_t$

#### Linear approximation

- $> x_{t+1} = h_t(x_t, \nu) = A_t(\nu)x_t + f_t(\nu)$ 
  - Where  $A_t(v) \in \mathbb{R}^{n \times n}$  is a matrix of linear function in terms of v
  - ► And  $f_t(v)$  is the difference (error) of  $x_{t+1}$  and  $A_t(v)x_t$
  - ► We assume this function to be a Gaussian process with certain kernel (e.g. radius based kernel)
  - So we can use Gaussian process regression to approximate this difference (error) function  $f_t(v) = x_{t+1} A_t(v)x_t$





$$A_1(\nu)$$
  $A_2(\nu)$  ...  $A_{T-1}(\nu)$   $A_{T-1}(\nu)$   $A_T(\nu)$   $A_T(\nu)$   $A_T(\nu)$   $A_T(\nu)$   $A_T(\nu)$   $A_T(\nu)$ 

$$\begin{split} x_k &= A_{k-1}(\nu) x_{k-1} + f_{k-1}(\nu) \\ &= A_{k-1}(\nu) \big( A_{k-2}(\nu) x_{k-2} + f_{k-2}(\nu) \big) + f_{k-1}(\nu) \\ &= A_{k-1}(\nu) A_{k-2}(\nu) x_{k-2} + A_{k-1}(\nu) f_{k-2}(\nu) + f_{k-1}(\nu) \\ &\vdots \\ &\vdots \\ &= p_1(\nu) f_1(\nu) + p_2(\nu) f_2(\nu) + \dots + p_{T-1}(\nu) f_{T-1}(\nu) + p_0(\nu) x_1 \end{split}$$

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$$x_{k} = A_{k-1}(\nu)x_{k-1} + f_{k-1}(\nu)$$

$$= A_{k-1}(\nu)(A_{k-2}(\nu)x_{k-2} + f_{k-2}(\nu)) + f_{k-1}(\nu)$$

$$= A_{k-1}(\nu)A_{k-2}(\nu)x_{k-2} + A_{k-1}(\nu)f_{k-2}(\nu) + f_{k-1}(\nu)$$
...
$$= p_{1}(\nu)f_{1}(\nu) + p_{2}(\nu)f_{2}(\nu) + \dots + p_{T-1}(\nu)f_{T-1}(\nu) + p_{0}(\nu)x_{1}$$

$$\sum x_{k} = g_{1}(\nu)f_{1}(\nu) + g_{2}(\nu)f_{2}(\nu) + \dots + g_{T-1}(\nu)f_{T-1}(\nu) + g_{T}(\nu)x_{1}$$

#### Kernel

- So we know the total infected population in the following T years would be
  - $f(v) = 1_I^T \sum x_k = g_1(v) 1_I^T f_1(v) + g_2(v) 1_I^T f_2(v) + \dots + g_{T-1}(v) 1_I^T f_{T-1}(v) + g_T(v) 1_I^T x_1$

#### Kernel

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  - $f(\nu) = 1_I^T \sum_{l} x_k = g_1(\nu) 1_I^T f_1(\nu) + g_2(\nu) 1_I^T f_2(\nu) + \dots + g_{T-1}(\nu) 1_I^T f_{T-1}(\nu) + g_T(\nu) 1_I^T x_1$
- Since we are assuming each simpler Gaussian process to have kernel  $k_i(\nu, \nu')$
- $\triangleright$  The entire kernel of function f is
  - $k(\nu, \nu') = g_1(\nu)k_1(\nu, \nu')g_1(\nu') + \dots + g_T(\nu)k_T(\nu, \nu')g_T(\nu')$

#### Gaussian process regression

- Now we can finally do the Gaussian process regression...
- $f(\nu) = g_1(\nu)f_1(\nu) + g_2(\nu)f_2(\nu) + \dots + g_T(\nu)f_T(\nu)$
- ► Then?

#### Gaussian process regression

- Now we have two options.
  - ▶ 1. run Gaussian process regression on the entire function f(v) with cumulative kernel k(v, v')
  - ▶ 2. run Gaussian process regression on each  $f_i(v)$  with kernel  $k_i(v, v')$  then sum them up.

#### Short story

- ▶ Long story short, the second one is better.
  - ▶ 1. run Gaussian process regression on the entire function f(v) with cumulative kernel k(v, v')
  - ▶ 2. run Gaussian process regression on each  $f_i(v)$  with kernel  $k_i(v, v')$  then sum them up.

#### Long story...

- Long story
  - ► We want to compare the variance derived from two different methods.
  - $var_{entire}(x) = k(x, x') k_T(x)^T K_T^{-1} k_T(x')$
  - $\triangleright var_i(x) = k_i(x, x') k_{i,T}(x)^T K_{i,T}^{-1} k_{i,T}(x')$

$$k_T(x, x') = k(x, x') - k_T(x)^T (K_T + \sigma^2 I)^{-1} k_T(x')$$
  
$$\sigma_T^2(x) = k_T(x, x)$$

# Long long story...

 $K_T = [k(x^j, x^k)]_{i,k} = \sum_i [g_i(x^j) k_i(x^j, x^k) g_i(x^k)]_{i,k} = \sum_i D_i K_{i,T} D_i$ 

 $D_i = diag([g_i(x^1), g_i(x^2), ..., g_i(x^m)])$ 

# Long long long story...

► 
$$k_{i,T}(x,x') = k_i(x,x') - k_{i,T}(x)^T K_{i,T}^{-1} k_{i,T}(x')$$
  
►  $var_i(x) = k_{i,T}(x,x)$   
►  $= k_i(x,x) - k_{i,T}(x)^T K_{i,T}^{-1} k_{i,T}(x)$   
►  $var(x) = \sum_i g_i(x) var_i(x) g_i(x)$   
►  $= \sum_i g_i(x) k_i(x,x) g_i(x) - \sum_i g_i(x) k_{i,T}(x)^T K_{i,T}^{-1} k_{i,T}(x) g_i(x)$   
►  $= \sum_i g_i(x) k_i(x,x) g_i(x) - \sum_i g_i(x) k_{i,T}(x)^T D_i D_i^{-1} K_{i,T}^{-1} D_i^{-1} D_i k_{i,T}(x) g_i(x)$   
►  $= \sum_i g_i(x) k_i(x,x) g_i(x) - \sum_i z_i^T D_i^{-1} K_{i,T}^{-1} D_i^{-1} z_i$   
►  $= \sum_i g_i(x) k_i(x,x) g_i(x) - \sum_i z_i^T B_i^{-1} z_i$ 

- $var_{entire}(x) = \sum_{i} g_i(x) k_i(x, x) g_i(x) \sum_{i,k} z_i^T \left(\sum_{j} D_j K_{j,T} D_j\right)^{-1} z_k$
- $= \sum_{i} g_i(x) k_i(x, x) g_i(x) \sum_{i,k} z_i^T \left(\sum_{j} B_j\right)^{-1} z_k$
- $var(x) = \sum_{i} g_i(x) k_i(x, x) g_i(x) \sum_{i} z_i^T B_i^{-1} z_i$
- $var_{entire}(x) var(x) = \sum_{i} z_i^T B_i^{-1} z_i \sum_{i,k} z_i^T \left(\sum_{j} B_j\right)^{-1} z_k$
- ▶ Where  $z_i \in R^m$ ,  $B_i \in S^m_+$  a positive semi definite m by m matrix.

#### ▶ We want to prove:

$$\Rightarrow \frac{\sum_{i} z_{i}^{T} B_{i}^{-1} z_{i}}{T} - \left(\frac{\sum_{i} z_{i}}{T}\right)^{T} \left(\frac{\sum_{j} B_{j}}{T}\right)^{-1} \left(\frac{\sum_{i} z_{i}}{T}\right) \geq 0$$

$$\Rightarrow \frac{\sum_{i} h(z_{i}, B_{i})}{T} - h\left(\frac{\sum_{i} z_{i}}{T}, \frac{\sum_{i} B_{i}}{T}\right) \geq 0$$

Where  $h(x, Y) = x^T Y^{-1} x$  is called a matrix fractional function, and is **convex** on  $dom \ h = \mathbf{R}^m \times S_+^m$ 

Therefore, by Jenson's inequality on convex function h, the inequality holds.

Therefore, by Jenson's inequality on convex function h, the inequality holds.

- ▶ i.e.  $var_{entire}(x) var(x) \ge 0$
- ▶ i.e.  $var_{entire}(x) \ge var(x)$

#### Conclusion

It implies running individual Gaussian process regression helps on reducing the uncertainty.

#### Next step

- ▶ Our problem is a two-stage problem:
  - ▶ 1. Learn the linear approximation  $g_i(v)$
  - ▶ 2. Approximate  $f(x) = \sum_i g_i(v) f_i(v)$  and find the optimal solution

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  - ▶ 1. Learn the linear approximation  $g_i(v)$
  - ▶ 2. Approximate  $f(x) = \sum_i g_i(v) f_i(v)$  and find the optimal solution
  - ▶ 3. Fit this method to our context...